



ENTERED

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/026,188
DATE: 07/15/2002
TIME: 13:55:39

Input Set : A:\Ucl149-1.app
Output Set: N:\CRF3\07152002\J026188.raw

```
4 <110> APPLICANT: Zuker, Charles S.
5     Zhang, Yifeng
6     The Regents of the University of California
8 <120> TITLE OF INVENTION: Assays for Taste Receptor Cell Specific
9     Ion Channel
11 <130> FILE REFERENCE: 02307E-114910US
13 <140> CURRENT APPLICATION NUMBER: US 10/026,188
14 <141> CURRENT FILING DATE: 2001-12-21
16 <150> PRIOR APPLICATION NUMBER: US 60/259,379
17 <151> PRIOR FILING DATE: 2000-12-29
19 <160> NUMBER OF SEQ ID NOS: 8
21 <170> SOFTWARE: FastSEQ for Windows Version 3.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 4185
25 <212> TYPE: DNA
26 <213> ORGANISM: Rattus sp.
28 <220> FEATURE:
29 <223> OTHER INFORMATION: rat L-TRP taste cDNA
31 <400> SEQUENCE: 1
32   caaaagcctc tggagagctg tgtcgagggt gttggaatcc agatgcccgg agttcgaaaa      60
33   gtcacaatgc cgatggccca gagctcttgt cctggaagcc cccagatac tggggatgga      120
34   tgggagccag tcctatgcaa gggagagggtc aacttcggag ggtctgggaa aaagcgaagc      180
35   aagtttgtga aggtgccaa gcaatgtggcc ccctccatgc tctttgaact cctgtctacc      240
36   gagtggcacc tgccagcccc caacctgggtg gtgtccctgg tgggcgagga acggcttttt      300
37   gctatgaagt cctggcttcg ggatgtcttg cgcaaggggc tggtgaaagc agctcagagc      360
38   acaggtgcct ggatcctgac cagtgccttc catgtgggcc tggcacgcca tgttggacag      420
39   gctgtacgtg atcactctct ggctagcaag tccaccaagg tccgtgtggt ggccatcgga      480
40   atggcctctc tggaccgaat ccttcaccgc caacttctag atggtgtcca ggaggatact      540
41   cccatccact acccagcaga tgaggggcag actcagggac cccctctgcc tctggacagc      600
42   aatctctccc acttcacct cgtggagcca ggcacccttg ggagtgggaa cgacggactg      660
43   gcagagctgc agctgagcct ggagaagcac atctctcagc agaggacagg ttatgggggt      720
44   accagcagca tccagatacc tgtcctttgc ttgctagtca atggtgacct cagcacccta      780
45   gagaggatgt ccagggcagt ggagcaggct gccccatggc tgatcctggc aggttctggg      840
46   ggcattgctg atgtactcgc tgccctgggt ggccagcctc atctcctggt gcccaggtg      900
47   accgagaagc agttcagaga gaaattccca agcgagtgtt tctcttgga agccattgta      960
48   cactggacag agctgctaca gaacattgct gcacaccccc acctgctcac agtgtacgac     1020
49   tttgagcagg agggttccga ggacctggac accgtcatcc tcaaggcact tgtgaaagcc     1080
50   tgcaagagtc acagccgaga cgcacaagac tacctagatg agctcaagtt agcagtggcc     1140
51   tgggatcgcg tggacattgc caagagtga atcttcaatg gggacgtgga gtggaagtcc     1200
52   tgtgacttgg aagagtgat gacagatgcc ctagttagca acaagcctga cttcgtgcgc     1260
53   ctctttgtgg acagtgtgac tgacatggcc gagttcttga cctatgggcg gctgcagcag     1320
54   ctttaccact ctgtgtcccc caagagcctc ctctttgaac tgctggagcg taagcatgag     1380
55   gagggtcggc tgacactggc tggcctgggt gcccagcaga cccggaagct gcccgttggt     1440
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/026,188

DATE: 07/15/2002

TIME: 13:55:39

Input Set : A:\Uc1149-1.app

Output Set: N:\CRF3\07152002\J026188.raw

```
56 ctgcctgcct tttcactcca tgagggtctcc cgagtttctca aagatttcct gcatgacgcc 1500
57 tgccgtggct tctaccagga tgggcgcagg atggagaaga gagggccacc caagcggcct 1560
58 gcaggccaga aatggctgcc ggacctcagt cggaagagtg aagacccatg gagggacctg 1620
59 ttcccttggg ctgtgctgca gaaccgttat gagatggcca catacttctg ggccatgggc 1680
60 cgggagggtg tggctgctgc tctggcggcc tgcaagatca tcaaggaaat gtccacctg 1740
61 gagaaagagg cagagggtggc ccgcactatg cgtgaggcca agtatgagca gctggccctc 1800
62 gatcttttct cagagtgtca cagcaacagt gaggacctg cctttgccct gttggtgcgc 1860
63 aggaaccaca gctggagcag gaccacctgc ctgcacctg ccactgaggc cgatgccaaag 1920
64 gccttctttg cccatgatgg tgtgcaagca ttcttgacga agatctggtg gggagacatg 1980
65 gccacaggca caccatctt acgacttctg ggtgccttca cctgcccagc cctcatctac 2040
66 acaaactctca tctccttcag tgaggatgcc ccgcagagga tggacctgga agatctgcag 2100
67 gagccagaca gtttgatata ggaaaagagc ttctgtgcca gccatggtgg ccaattggag 2160
68 aagttaacag aggcgcacaag ggctcctggc gatctaggcc cacaagctgc cttcctgctc 2220
69 acacggtgga ggaagtctct gggcgctcct gtgactgtgt tcttggggaa tgtggtcatg 2280
70 tactttgcat tctcttccct atttctctac gtccctgctg tggatttcag gccaccaccc 2340
71 caggggccat ctgggtcgga agttaccctg tatttctggg tcttcacact ggtgctggag 2400
72 gaaatccgac agggattctt cacaacagag gacaccgctc tggtaagaa gttcactctg 2460
73 tacgtagaag acaactggaa caaatgtgac atggtggcca tcttctgtt cattgttgg 2520
74 gtcacctgta gaatggtgcc ctccgtgttt gaggtggcc ggaactgttct ggccattgac 2580
75 ttcattggtg tccacttcg gctcatccac atctttgcta ttcacaagca gctgggtcct 2640
76 aagatcatca ttgtagagcg gatgatgaaa gatgtcttct tcttctctt cttcctgagc 2700
77 gtgtggctcg tggcctatgg cgtgaccact caggccctgc tggaccccca cgatggccgt 2760
78 ctggagtgga ttttcgcgcg tgtgctctac aggccttacc tgcagatctt tgggcaaatc 2820
79 cctctggatg aaattgatga ggcccggtg aactgctctc ttcaccggtt gctgctggac 2880
80 agctcagctt cctgccctaa tctctatgcc aactggctgg tcattctcct gctggttacc 2940
81 ttctctctcg tactaatgt gctacttatg aaccttctga tcgccatgtt cagctacaca 3000
82 ttccaggtgg tgcagggcaa tgcagacatg ttctggaagt ttcaacgcta ccacctcatc 3060
83 gttgaatacc acggaaggcc ggctctggcc ccgcccttca tctgctcag ccacctgagc 3120
84 ctggtgctca agcaggctct caggaaggaa gccagcaca aacagcaaca cctggagaga 3180
85 gacttgctg acccctggtg ccagaagatc attacctggg aaacagttca aaaggagaac 3240
86 ttctgagta ccatggagaa acggaggagg gacagtgaga aggaggtgct gaggaaaacg 3300
87 gcacacagag tggacttgat tgccaaatac atcgggggtc tgagagagca agaaaagagg 3360
88 atcaagtgtc tggagtccca ggccaactac tgtatgtctc tcttgtcctc catgactgac 3420
89 aactggctc ctggaggcac ctactcaagt tctcaaaact gtggtcgag ggtcagcca 3480
90 gcctctgcta gagacagga gtacctagag gctggcttgc cactcaga cactgaaat 3540
91 ggagaaacca cttgccctag agctccagac ctggccagat tgaggttttg ggtcacatca 3600
92 accttccct gcccccagca gcccagagac cttgccgag accatgtctt ggacacctct 3660
93 tcctatgaaa atgagactca tgtctttggc atctatctgg gagccccagg cgtcctctcc 3720
94 agcaggggaa gttttctcat gtcctacctc aaactttcac cagctaagac tggacagctg 3780
95 gaactggcca agtcccatat gggataccat ctgcctggat ggggctactt acgtctagcc 3840
96 tgtcttacc tgagttccaa agaggccaac ctcttaaaca ctagaggttt cttcttgtc 3900
97 ctctgatcca tccatcagcc gaccagcttc tagaggcgag gactcagatc tactgtaatc 3960
98 agctcccatc cttcagcccc cacagcataa tttgtgtgat tgtcctggca caaaccccaa 4020
99 gatactgctc aagggtaccc aatgctatct tactttctat aaagcctgta gaccacctca 4080
100 actaagctaa actggaccac aggggtggct aaaccaacat ttcaaacacc tggggaacat 4140
101 ggagttatct gacccaaaaa aaaaaaaaaa aaaaaaaaaa aaaaa 4185
103 <210> SEQ ID NO: 2
104 <211> LENGTH: 1165
105 <212> TYPE: PRT
```

RAW SEQUENCE LISTING

DATE: 07/15/2002

PATENT APPLICATION: US/10/026,188

TIME: 13:55:39

Input Set : A:\Uc1149-1.app

Output Set: N:\CRF3\07152002\J026188.raw

106 <213> ORGANISM: Rattus sp.
108 <220> FEATURE:
109 <223> OTHER INFORMATION: rat L-TRP taste predicted protein
111 <400> SEQUENCE: 2

112	Met	Pro	Gly	Val	Arg	Lys	Val	Thr	Met	Pro	Met	Ala	Gln	Ser	Ser	Cys
113	1				5				10					15		
114	Pro	Gly	Ser	Pro	Pro	Asp	Thr	Gly	Asp	Gly	Trp	Glu	Pro	Val	Leu	Cys
115				20					25					30		
116	Lys	Gly	Glu	Val	Asn	Phe	Gly	Gly	Ser	Gly	Lys	Lys	Arg	Ser	Lys	Phe
117				35				40					45			
118	Val	Lys	Val	Pro	Ser	Asn	Val	Ala	Pro	Ser	Met	Leu	Phe	Glu	Leu	Leu
119		50					55					60				
120	Leu	Thr	Glu	Trp	His	Leu	Pro	Ala	Pro	Asn	Leu	Val	Val	Ser	Leu	Val
121	65					70					75					80
122	Gly	Glu	Glu	Arg	Leu	Phe	Ala	Met	Lys	Ser	Trp	Leu	Arg	Asp	Val	Leu
123					85					90					95	
124	Arg	Lys	Gly	Leu	Val	Lys	Ala	Ala	Gln	Ser	Thr	Gly	Ala	Trp	Ile	Leu
125				100					105					110		
126	Thr	Ser	Ala	Leu	His	Val	Gly	Leu	Ala	Arg	His	Val	Gly	Gln	Ala	Val
127				115				120					125			
128	Arg	Asp	His	Ser	Leu	Ala	Ser	Thr	Ser	Thr	Lys	Val	Arg	Val	Val	Ala
129		130					135					140				
130	Ile	Gly	Met	Ala	Ser	Leu	Asp	Arg	Ile	Leu	His	Arg	Gln	Leu	Leu	Asp
131	145					150					155					160
132	Gly	Val	Gln	Glu	Asp	Thr	Pro	Ile	His	Tyr	Pro	Ala	Asp	Glu	Gly	Ser
133					165					170					175	
134	Thr	Gln	Gly	Pro	Leu	Cys	Pro	Leu	Asp	Ser	Asn	Leu	Ser	His	Phe	Ile
135				180					185					190		
136	Leu	Val	Glu	Pro	Gly	Thr	Leu	Gly	Ser	Gly	Asn	Asp	Gly	Leu	Ala	Glu
137			195					200					205			
138	Leu	Gln	Leu	Ser	Leu	Glu	Lys	His	Ile	Ser	Gln	Gln	Arg	Thr	Gly	Tyr
139		210					215						220			
140	Gly	Gly	Thr	Ser	Ser	Ile	Gln	Ile	Pro	Val	Leu	Cys	Leu	Leu	Val	Asn
141	225					230					235					240
142	Gly	Asp	Pro	Ser	Thr	Leu	Glu	Arg	Met	Ser	Arg	Ala	Val	Glu	Gln	Ala
143					245					250					255	
144	Ala	Pro	Trp	Leu	Ile	Leu	Ala	Gly	Ser	Gly	Gly	Ile	Ala	Asp	Val	Leu
145				260					265					270		
146	Ala	Ala	Leu	Val	Gly	Gln	Pro	His	Leu	Leu	Val	Pro	Gln	Val	Thr	Glu
147				275					280					285		
148	Lys	Gln	Phe	Arg	Glu	Lys	Phe	Pro	Ser	Glu	Cys	Phe	Ser	Trp	Glu	Ala
149		290					295						300			
150	Ile	Val	His	Trp	Thr	Glu	Leu	Leu	Gln	Asn	Ile	Ala	Ala	His	Pro	His
151	305					310					315					320
152	Leu	Leu	Thr	Val	Tyr	Asp	Phe	Glu	Gln	Glu	Gly	Ser	Glu	Asp	Leu	Asp
153					325					330					335	
154	Thr	Val	Ile	Leu	Lys	Ala	Leu	Val	Lys	Ala	Cys	Lys	Ser	His	Ser	Arg
155				340					345					350		
156	Asp	Ala	Gln	Asp	Tyr	Leu	Asp	Glu	Leu	Lys	Leu	Ala	Val	Ala	Trp	Asp

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/026,188

DATE: 07/15/2002

TIME: 13:55:39

Input Set : A:\Uc1149-1.app

Output Set: N:\CRF3\07152002\J026188.raw

157					355											360															365
158	Arg	Val	Asp	Ile	Ala	Lys	Ser	Glu	Ile	Phe	Asn	Gly	Asp	Val	Glu	Trp															
159						370											375														380
160	Lys	Ser	Cys	Asp	Leu	Glu	Glu	Val	Met	Thr	Asp	Ala	Leu	Val	Ser	Asn															
161						385											390														400
162	Lys	Pro	Asp	Phe	Val	Arg	Leu	Phe	Val	Asp	Ser	Gly	Ala	Asp	Met	Ala															
163						405											410														415
164	Glu	Phe	Leu	Thr	Tyr	Gly	Arg	Leu	Gln	Gln	Leu	Tyr	His	Ser	Val	Ser															
165						420											425														430
166	Pro	Lys	Ser	Leu	Leu	Phe	Glu	Leu	Leu	Glu	Arg	Lys	His	Glu	Glu	Gly															
167						435											440														445
168	Arg	Leu	Thr	Leu	Ala	Gly	Leu	Gly	Ala	Gln	Gln	Thr	Arg	Lys	Leu	Pro															
169						450											455														460
170	Val	Gly	Leu	Pro	Ala	Phe	Ser	Leu	His	Glu	Val	Ser	Arg	Val	Leu	Lys															
171						465											470														480
172	Asp	Phe	Leu	His	Asp	Ala	Cys	Arg	Gly	Phe	Tyr	Gln	Asp	Gly	Arg	Arg															
173						485											490														495
174	Met	Glu	Lys	Arg	Gly	Pro	Pro	Lys	Arg	Pro	Ala	Gly	Gln	Lys	Trp	Leu															
175						500											505														510
176	Pro	Asp	Leu	Ser	Arg	Lys	Ser	Glu	Asp	Pro	Trp	Arg	Asp	Leu	Phe	Leu															
177						515											520														525
178	Trp	Ala	Val	Leu	Gln	Asn	Arg	Tyr	Glu	Met	Ala	Thr	Tyr	Phe	Trp	Ala															
179						530											535														540
180	Met	Gly	Arg	Glu	Gly	Val	Ala	Ala	Ala	Leu	Ala	Ala	Cys	Lys	Ile	Ile															
181						545											550														555
182	Lys	Glu	Met	Ser	His	Leu	Glu	Lys	Glu	Ala	Glu	Val	Ala	Arg	Thr	Met															
183						565											570														575
184	Arg	Glu	Ala	Lys	Tyr	Glu	Gln	Leu	Ala	Leu	Asp	Leu	Phe	Ser	Glu	Cys															
185						580											585														590
186	Tyr	Ser	Asn	Ser	Glu	Asp	Arg	Ala	Phe	Ala	Leu	Leu	Val	Arg	Arg	Asn															
187						595											600														605
188	His	Ser	Trp	Ser	Arg	Thr	Thr	Cys	Leu	His	Leu	Ala	Thr	Glu	Ala	Asp															
189						610											615														620
190	Ala	Lys	Ala	Phe	Phe	Ala	His	Asp	Gly	Val	Gln	Ala	Phe	Leu	Thr	Lys															
191						625											630														635
192	Ile	Trp	Trp	Gly	Asp	Met	Ala	Thr	Gly	Thr	Pro	Ile	Leu	Arg	Leu	Leu															
193						645											650														655
194	Gly	Ala	Phe	Thr	Cys	Pro	Ala	Leu	Ile	Tyr	Thr	Asn	Leu	Ile	Ser	Phe															
195						660											665														670
196	Ser	Glu	Asp	Ala	Pro	Gln	Arg	Met	Asp	Leu	Glu	Asp	Leu	Gln	Glu	Pro															
197						675											680														685
198	Asp	Ser	Leu	Asp	Met	Glu	Lys	Ser	Phe	Leu	Cys	Ser	His	Gly	Gly	Gln															
199						690											695														700
200	Leu	Glu	Lys	Leu	Thr	Glu	Ala	Pro	Arg	Ala	Pro	Gly	Asp	Leu	Gly	Pro															
201						705											710														715
202	Gln	Ala	Ala	Phe	Leu	Thr	Arg	Trp	Arg	Lys	Phe	Trp	Gly	Ala	Pro																
203						725											730														735
204	Val	Thr	Val	Phe	Leu	Gly	Asn	Val	Val	Met	Tyr	Phe	Ala	Phe	Leu	Phe															
205						740											745														750

RAW SEQUENCE LISTING

DATE: 07/15/2002

PATENT APPLICATION: US/10/026,188

TIME: 13:55:39

Input Set : A:\Uc1149-1.app

Output Set: N:\CRF3\07152002\J026188.raw

206	Leu	Phe	Ser	Tyr	Val	Leu	Leu	Val	Asp	Phe	Arg	Pro	Pro	Pro	Gln	Gly
207			755					760					765			
208	Pro	Ser	Gly	Ser	Glu	Val	Thr	Leu	Tyr	Phe	Trp	Val	Phe	Thr	Leu	Val
209			770				775					780				
210	Leu	Glu	Glu	Ile	Arg	Gln	Gly	Phe	Phe	Thr	Asn	Glu	Asp	Thr	Arg	Leu
211			785			790					795					800
212	Val	Lys	Lys	Phe	Thr	Leu	Tyr	Val	Glu	Asp	Asn	Trp	Asn	Lys	Cys	Asp
213				805						810					815	
214	Met	Val	Ala	Ile	Phe	Leu	Phe	Ile	Val	Gly	Val	Thr	Cys	Arg	Met	Val
215				820						825				830		
216	Pro	Ser	Val	Phe	Glu	Ala	Gly	Arg	Thr	Val	Leu	Ala	Ile	Asp	Phe	Met
217			835						840					845		
218	Val	Phe	Thr	Leu	Arg	Leu	Ile	His	Ile	Phe	Ala	Ile	His	Lys	Gln	Leu
219		850					855					860				
220	Gly	Pro	Lys	Ile	Ile	Ile	Val	Glu	Arg	Met	Met	Lys	Asp	Val	Phe	Phe
221		865				870					875					880
222	Phe	Leu	Phe	Phe	Leu	Ser	Val	Trp	Leu	Val	Ala	Tyr	Gly	Val	Thr	Thr
223				885						890					895	
224	Gln	Ala	Leu	Leu	Asp	Pro	His	Asp	Gly	Arg	Leu	Glu	Trp	Ile	Phe	Arg
225				900					905					910		
226	Arg	Val	Leu	Tyr	Arg	Pro	Tyr	Leu	Gln	Ile	Phe	Gly	Gln	Ile	Pro	Leu
227			915					920					925			
228	Asp	Glu	Ile	Asp	Glu	Ala	Arg	Val	Asn	Cys	Ser	Leu	His	Pro	Leu	Leu
229		930					935					940				
230	Leu	Asp	Ser	Ser	Ala	Ser	Cys	Pro	Asn	Leu	Tyr	Ala	Asn	Trp	Leu	Val
231		945				950					955					960
232	Ile	Leu	Leu	Leu	Val	Thr	Phe	Leu	Leu	Val	Thr	Asn	Val	Leu	Leu	Met
233				965						970					975	
234	Asn	Leu	Leu	Ile	Ala	Met	Phe	Ser	Tyr	Thr	Phe	Gln	Val	Val	Gln	Gly
235				980					985					990		
236	Asn	Ala	Asp	Met	Phe	Trp	Lys	Phe	Gln	Arg	Tyr	His	Leu	Ile	Val	Glu
237		995					1000						1005			
238	Tyr	His	Gly	Arg	Pro	Ala	Leu	Ala	Pro	Pro	Phe	Ile	Leu	Leu	Ser	His
239		1010					1015					1020				
240	Leu	Ser	Leu	Val	Leu	Lys	Gln	Val	Phe	Arg	Lys	Glu	Ala	Gln	His	Lys
241		1025				1030					1035					1040
242	Gln	Gln	His	Leu	Glu	Arg	Asp	Leu	Pro	Asp	Pro	Val	Asp	Gln	Lys	Ile
243				1045						1050					1055	
244	Ile	Thr	Trp	Glu	Thr	Val	Gln	Lys	Glu	Asn	Phe	Leu	Ser	Thr	Met	Glu
245			1060						1065					1070		
246	Lys	Arg	Arg	Arg	Asp	Ser	Glu	Lys	Glu	Val	Leu	Arg	Lys	Thr	Ala	His
247			1075					1080					1085			
248	Arg	Val	Asp	Leu	Ile	Ala	Lys	Tyr	Ile	Gly	Gly	Leu	Arg	Glu	Gln	Glu
249		1090					1095					1100				
250	Lys	Arg	Ile	Lys	Cys	Leu	Glu	Ser	Gln	Ala	Asn	Tyr	Cys	Met	Leu	Leu
251		1105				1110					1115					1120
252	Leu	Ser	Ser	Met	Thr	Asp	Thr	Leu	Ala	Pro	Gly	Gly	Thr	Tyr	Ser	Ser
253				1125						1130					1135	
254	Ser	Gln	Asn	Cys	Gly	Arg	Arg	Ser	Gln	Pro	Ala	Ser	Ala	Arg	Asp	Arg

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/026,188

DATE: 07/15/2002

TIME: 13:55:40

Input Set : A:\Uc1149-1.app

Output Set: N:\CRF3\07152002\J026188.raw